PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/031,014

DATE: 02/07/2002
TIME: 16:15:28

Input Set : A:\20493p.txt
Output Set: N:\CRF3\02072002\J031014.raw

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      4 <110> APPLICANT: Howard, Andrew D.
      6 <120> TITLE OF INVENTION: Melanin-Concentrating Hormone Receptor
      9 <130> FILE REFERENCE: 20493P
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/031,014
C--> 11 <141> CURRENT FILING DATE: 2002-01-10
     11 <150> PRIOR APPLICATION NUMBER: PCT/US00/18733
     12 <151> PRIOR FILING DATE: 2000-07-10
     14 <150> PRIOR APPLICATION NUMBER: US 60/143,706
     15 <151> PRIOR FILING DATE: 1999-07-14
     17 <160> NUMBER OF SEQ ID NOS: 10
     19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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     29 atcatcatgc cttcggtgtt cggcaccatc tgcctcctgg gcatcatcgg gaactccacg   180
     30 gtcatcttcg cggtcgtgaa gaagtccaag ctgcactggt gcaacaacgt ccccgacatc   240
     31 ttcatcatca acctctcggt agtagatctc ctctttctcc tgggcatgcc cttcatgatc   300
     32 caccagctca tgggcaatgg ggtgtggcac tttggggaga ccatgtgcac cctcatcacg   360
     33 gccatggatg ccaatagtca gttcaccagc acctacatcc tgaccgccat ggccattgac   420
     34 cgctacctgg ccactgtcca ccccatctct tccacgaagt tccggaagcc ctctgtggcc   480
     35 accctggtga tctgcctcct gtgggccctc tccttcatca gcatcacccc tgtgtggctg   540
     36 tatgccagac tcatcccctt cccaggaggt gcagtgggct gcggcatacg cctgcccaac   600
     37 ccagacactg acctctactg gttcaccctg taccagtttt tcctggcctt tgccctgcct   660
     38 tttgtggtca tcacagccgc atacgtgagg atcctgcagc gcatgacgtc ctcagtggcc   720
     39 cccgcctccc agcgcagcat ccggctgcgg acaaagaggg tgacccgcac agccatcgcc   780
     40 atctgtctgg tcttctttgt gtgctgggca ccctactatg tgctacagct gacccagttg   840
     41 tccatcagcc gcccgaccct cacctttgtc tacttataca atgcggccat cagcttgggc   900
     42 tatgccaaca gctgcctcaa cccctttgtg tacatcgtgc tctgtgagac gttccgcaaa   960
     43 cgcttggtcc tgtcggtgaa gcctgcagcc caggggcagc ttcgcgctgt cagcaacgct  1020
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     47 <211> LENGTH: 353
     48 <212> TYPE: PRT
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     51 <400> SEQUENCE: 2
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     53  1               5                  10                  15
     54 Thr Ser Asp Gly Pro Asp Asn Leu Thr Ser Ala Gly Ser Pro Pro Arg
     55              20                  25                  30
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56 Thr Gly Ser Ile Ser Tyr Ile Asn Ile Ile Met Pro Ser Val Phe Gly
57         35              40              45
58 Thr Ile Cys Leu Leu Gly Ile Ile Gly Asn Ser Thr Val Ile Phe Ala
59     50              55              60
60 Val Val Lys Lys Ser Lys Leu His Trp Cys Asn Asn Val Pro Asp Ile
61 65              70              75              80
62 Phe Ile Ile Asn Leu Ser Val Val Asp Leu Leu Phe Leu Leu Gly Met
63             85              90              95
64 Pro Phe Met Ile His Gln Leu Met Gly Asn Gly Val Trp His Phe Gly
65         100             105             110
66 Glu Thr Met Cys Thr Leu Ile Thr Ala Met Asp Ala Asn Ser Gln Phe
67         115             120             125
68 Thr Ser Thr Tyr Ile Leu Thr Ala Met Ala Ile Asp Arg Tyr Leu Ala
69     130             135             140
70 Thr Val His Pro Ile Ser Ser Thr Lys Phe Arg Lys Pro Ser Val Ala
71 145             150             155             160
72 Thr Leu Val Ile Cys Leu Leu Trp Ala Leu Ser Phe Ile Ser Ile Thr
73             165             170             175
74 Pro Val Trp Leu Tyr Ala Arg Leu Ile Pro Phe Pro Gly Gly Ala Val
75             180             185             190
76 Gly Cys Gly Ile Arg Leu Pro Asn Pro Asp Thr Asp Leu Tyr Trp Phe
77         195             200             205
78 Thr Leu Tyr Gln Phe Phe Leu Ala Phe Ala Leu Pro Phe Val Val Ile
79     210             215             220
80 Thr Ala Ala Tyr Val Arg Ile Leu Gln Arg Met Thr Ser Ser Val Ala
81 225             230             235             240
82 Pro Ala Ser Gln Arg Ser Ile Arg Leu Arg Thr Lys Arg Val Thr Arg
83             245             250             255
84 Thr Ala Ile Ala Ile Cys Leu Val Phe Phe Val Cys Trp Ala Pro Tyr
85         260             265             270
86 Tyr Val Leu Gln Leu Thr Gln Leu Ser Ile Ser Arg Pro Thr Leu Thr
87         275             280             285
88 Phe Val Tyr Leu Tyr Asn Ala Ala Ile Ser Leu Gly Tyr Ala Asn Ser
89     290             295             300
90 Cys Leu Asn Pro Phe Val Tyr Ile Val Leu Cys Glu Thr Phe Arg Lys
91 305             310             315             320
92 Arg Leu Val Leu Ser Val Lys Pro Ala Ala Gln Gly Gln Leu Arg Ala
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96 Thr
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101 <212> TYPE: DNA
102 <213> ORGANISM: Human
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107 aggctgccgc agcctgcgtg ggtggagggg agctcagctc ggttgtggga gcaggcgacc    180
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109 acctctgatg gccccgataa cctcacttcg gcaggatcac ctcctcgcac ggggagcatc   300
110 tcctacatca acatcatcat gccttcggtg ttcggcacca tctgcctcct gggcatcatc   360
111 gggaactcca cggtcatctt cgcggtcgtg aagaagtcca agctgcactg gtgcaacaac   420
112 gtccccgaca tcttcatcat caacctctcg gtagtagatc tcctctttct cctgggcatg   480
113 cccttcatga tccaccagct catgggcaat ggggtgtggc actttgggga gaccatgtgc   540
114 accctcatca cggccatgga tgccaatagt cagttcacca gcacctacat cctgaccgcc   600
115 atggccattg accgctacct ggccactgtc caccccatct cttccacgaa gttccggaag   660
116 ccctctgtgg ccaccctggt gatctgcctc ctgtgggccc tctccttcat cagcatcacc   720
117 cctgtgtggc tgtatgccag actcatcccc ttcccaggag gtgcagtggg ctgcggcata   780
118 cgcctgccca acccagacac tgacctctac tggttcaccc tgtaccagtt tttcctggcc   840
119 tttgccctgc cttttgtggt catcacagcc gcatacgtga ggatcctgca gcgcatgacg   900
120 tcctcagtgg cccccgcctc ccagcgcagc atccggctgc ggacaaagag ggtgacccgc   960
121 acagccatcg ccatctgtct ggtcttcttt gtgtgctggg caccctacta tgtgctacag  1020
122 ctgacccagt tgtccatcag ccgcccgacc ctcacctttg tctacttata caatgcggcc  1080
123 atcagcttgg gctatgccaa cagctgcctc aacccctttg tgtacatcgt gctctgtgag  1140
124 acgttccgca aacgcttggt cctgtcggtg aagcctgcag cccagggggca gcttcgcgct  1200
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128 <211> LENGTH: 417
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136             20                  25                  30
137 Pro Gly Gln Gly Gly Arg Arg Trp Arg Leu Pro Gln Pro Ala Trp Val
138          35                  40                  45
139 Glu Gly Ser Ser Ala Arg Leu Trp Glu Gln Ala Thr Gly Thr Gly Trp
140       50                  55                  60
141 Met Asp Leu Glu Ala Ser Leu Leu Pro Thr Gly Pro Asn Ala Ser Asn
142 65                  70                  75                  80
143 Thr Ser Asp Gly Pro Asp Asn Leu Thr Ser Ala Gly Ser Pro Pro Arg
144             85                  90                  95
145 Thr Gly Ser Ile Ser Tyr Ile Asn Ile Ile Met Pro Ser Val Phe Gly
146             100                 105                 110
147 Thr Ile Cys Leu Leu Gly Ile Ile Gly Asn Ser Thr Val Ile Phe Ala
148          115                 120                 125
149 Val Val Lys Lys Ser Lys Leu His Trp Cys Asn Asn Val Pro Asp Ile
150       130                 135                 140
151 Phe Ile Ile Asn Leu Ser Val Val Asp Leu Leu Phe Leu Leu Gly Met
152 145                 150                 155                 160
153 Pro Phe Met Ile His Gln Leu Met Gly Asn Gly Val Trp His Phe Gly
154             165                 170                 175
155 Glu Thr Met Cys Thr Leu Ile Thr Ala Met Asp Ala Asn Ser Gln Phe
156             180                 185                 190
157 Thr Ser Thr Tyr Ile Leu Thr Ala Met Ala Ile Asp Arg Tyr Leu Ala
158             195                 200                 205
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159 Thr Val His Pro Ile Ser Ser Thr Lys Phe Arg Lys Pro Ser Val Ala
160     210             215             220
161 Thr Leu Val Ile Cys Leu Leu Trp Ala Leu Ser Phe Ile Ser Ile Thr
162 225             230             235             240
163 Pro Val Trp Leu Tyr Ala Arg Leu Ile Pro Phe Pro Gly Gly Ala Val
164             245             250             255
165 Gly Cys Gly Ile Arg Leu Pro Asn Pro Asp Thr Asp Leu Tyr Trp Phe
166             260             265             270
167 Thr Leu Tyr Gln Phe Phe Leu Ala Phe Ala Leu Pro Phe Val Val Ile
168     275             280             285
169 Thr Ala Ala Tyr Val Arg Ile Leu Gln Arg Met Thr Ser Ser Val Ala
170     290             295             300
171 Pro Ala Ser Gln Arg Ser Ile Arg Leu Arg Thr Lys Arg Val Thr Arg
172 305             310             315             320
173 Thr Ala Ile Ala Ile Cys Leu Val Phe Phe Val Cys Trp Ala Pro Tyr
174             325             330             335
175 Tyr Val Leu Gln Leu Thr Gln Leu Ser Ile Ser Arg Pro Thr Leu Thr
176             340             345             350
177 Phe Val Tyr Leu Tyr Asn Ala Ala Ile Ser Leu Gly Tyr Ala Asn Ser
178     355             360             365
179 Cys Leu Asn Pro Phe Val Tyr Ile Val Leu Cys Glu Thr Phe Arg Lys
180     370             375             380
181 Arg Leu Val Leu Ser Val Lys Pro Ala Ala Gln Gly Gln Leu Arg Ala
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191 <213> ORGANISM: Human
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196 ggtggcaggc gctggaggct gccgcagcct gcgtgggtgg aggggagctc agctcggttg     180
197 tgggagcagg cgaccggcac tggctggatg gacctggaag cctcgctgct gcccactggt     240
198 cccaacgcca gcaacacctc tgatggcccc gataacctca cttcggcagg atcacctcct     300
199 cgcacgggga gcatctccta catcaacatc atcatgcctt cggtgttcgg caccatctgc     360
200 ctcctgggca tcatcgggaa ctccacggtc atcttcgcgg tcgtgaagaa gtccaagctg     420
201 cactggtgca acaacgtccc cgacatcttc atcatcaacc tctcggtagt agatctcctc     480
202 tttctcctgg gcatgccctt catgatccac cagctcatgg gcaatggggt gtggcacttt     540
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207 gtgggctgcg gcatacgcct gcccaaccca gacactgacc tctactggtt caccctgtac     840
208 cagttttttcc tggcctttgc cctgccttttt gtggtcatca cagccgcata cgtgaggatc     900
209 ctgcagcgca tgacgtcctc agtggccccc gcctcccagc gcagcatccg gctgcggaca     960
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Input Set : A:\20493p.txt
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212 ttatacaatg cggccatcag cttgggctat gccaacagct gcctcaaccc ctttgtgtac   1140
213 atcgtgctct gtgagacgtt ccgcaaacgc ttggtcctgt cggtgaagcc tgcagcccag   1200
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227 Cys Gly Ala Cys Ala Pro Gly Gln Gly Gly Arg Arg Trp Arg Leu Pro
228         35                  40                  45
229 Gln Pro Ala Trp Val Glu Gly Ser Ser Ala Arg Leu Trp Glu Gln Ala
230     50                  55                  60
231 Thr Gly Thr Gly Trp Met Asp Leu Glu Ala Ser Leu Leu Pro Thr Gly
232 65                  70                  75                  80
233 Pro Asn Ala Ser Asn Thr Ser Asp Gly Pro Asp Asn Leu Thr Ser Ala
234                 85                  90                  95
235 Gly Ser Pro Pro Arg Thr Gly Ser Ile Ser Tyr Ile Asn Ile Ile Met
236             100                 105                 110
237 Pro Ser Val Phe Gly Thr Ile Cys Leu Leu Gly Ile Ile Gly Asn Ser
238             115                 120                 125
239 Thr Val Ile Phe Ala Val Val Lys Lys Ser Lys Leu His Trp Cys Asn
240         130                 135                 140
241 Asn Val Pro Asp Ile Phe Ile Ile Asn Leu Ser Val Val Asp Leu Leu
242 145                 150                 155                 160
243 Phe Leu Leu Gly Met Pro Phe Met Ile His Gln Leu Met Gly Asn Gly
244                 165                 170                 175
245 Val Trp His Phe Gly Glu Thr Met Cys Thr Leu Ile Thr Ala Met Asp
246             180                 185                 190
247 Ala Asn Ser Gln Phe Thr Ser Thr Tyr Ile Leu Thr Ala Met Ala Ile
248         195                 200                 205
249 Asp Arg Tyr Leu Ala Thr Val His Pro Ile Ser Ser Thr Lys Phe Arg
250     210                 215                 220
251 Lys Pro Ser Val Ala Thr Leu Val Ile Cys Leu Leu Trp Ala Leu Ser
252 225                 230                 235                 240
253 Phe Ile Ser Ile Thr Pro Val Trp Leu Tyr Ala Arg Leu Ile Pro Phe
254                 245                 250                 255
255 Pro Gly Gly Ala Val Gly Cys Gly Ile Arg Leu Pro Asn Pro Asp Thr
256             260                 265                 270
257 Asp Leu Tyr Trp Phe Thr Leu Tyr Gln Phe Phe Leu Ala Phe Ala Leu
258             275                 280                 285
259 Pro Phe Val Val Ile Thr Ala Ala Tyr Val Arg Ile Leu Gln Arg Met
260     290                 295                 300
261 Thr Ser Ser Val Ala Pro Ala Ser Gln Arg Ser Ile Arg Leu Arg Thr
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VERIFICATION SUMMARY   DATE: 02/07/2002
PATENT APPLICATION: US/10/031,014   TIME: 16:15:29

Input Set : A:\20493p.txt
Output Set: N:\CRF3\02072002\J031014.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date